Figure 2

Mutation	Exemplary Pool	3' blocking group	Oligo type (Arm)	Sequence (5'-3')	ε_{260} M^{-1} cm^{-1}
2789+5G>A	1	none	invader	TTTGGTTGTGCTGTGGCTCCTTGGAAAGTGAT (SEQ ID NO:1)	330800
2789+5G>A	1	hex	probe/DM	CGCGCCGAGGATATTCCATGTCCTATTGTG (SEQ ID NO:2)	306500
2789+5G>A	1	none	synthetic target	CAATCTACACAATAGGACATGGAATATTCACTTTCCAAGGAGCCACAGCACAACCAAA (SEQ ID NO:3)	667000
R1162X	1	none	invader	GTTTACCTTCTGTTGGCATGTCAATGAACTTAAAGACTCT (SEQ ID NO:4)	428000
R1162X	1	hex	probe/DM	CGCGCCGAGGAGCTCACAGATCGC (SEQ ID NO:5)	253000
R1162X	1	none	synthetic target	TCAGATGCGATCTGTGAGCTGAGTCTTTAAGTTCATTGACATGCCAACAGAAGGTAAAC (SEQ ID NO:6)	659000
R347P	4	none	invader	CAGGGAAATTGCCGAGTGACCGCCATGT (SEQ ID NO:7)	306600
R347P	4	hex	probe/ER24	ACGGACGCGGAGGGCAGAACAATGCAG (SEQ ID NO:8)	318200
R347P	4	none	synthetic target	CTCATTCTGCATTGTTCTGCCCATGGCGGTCACTCGGCAATTTCCCTGGG (SEQ ID NO:9)	488000
1898+1G>A	1	none	invader	GACTCTCCTTTTGGATACCTAGATGTTTTAACAGAAAAAGAAATATTTGAAAGT (SEQ ID NO:10)	619900
1898+1G>A	1	hex	probe/DM	CGCGCCGAGGATATGTTCTTTGAATACCTTACTTAT (SEQ ID NO:11)	386500
1898+1G>A	1	none	synthetic target	ATAAGTAAGGTATTCAAAGAACATATCTTTCAAATATTTCTTTTTCTGTTAAAACATCTAGGTATCCAAAAGGAGAGTC (SEQ ID NO:12)	904800
2184delA	4	none	invader	CCCCAAACTCTCCAGTCTGTTTAAAAGATTATTTTTTC (SEQ ID NO:13)	393000
2184delA	4	hex	probe/DM	CGCGCCGAGGGTTTCTGTCCAGGAGACA (SEQ ID NO:14)	305200
del1507	1	none	invader	GCTTTGATGACGCTTCTGTATCTATATTCATCATAGGAAACACCAAT (SEQ ID NO:15)	509300
del1507	1	hex	probe/DM	CGCGCCGAGGAGATATTTTCTTTAATGGTGCC (SEQ ID NO:16)	345200
del1507	1	none	synthetic target	GCCTGGCACCATTAAAGAAAATATCTTTGGTGTTTCCTATGATGAATATAGATACAGAAGCGTCATCAAAGCATGCC (SEQ ID NO:17)	866600
G85E	4	none	invader	GCCCTTCGGCGATGTTTTTTCTGGAGATTTATGTTCTATGT (SEQ ID NO:18)	409100
G85E	4	hex	probe/ER24	ACGGACGCGGAGAAATCTTTTTATATTTAGGGGTAAG (SEQ ID NO:19)	431700
G85E	4	none	synthetic target	AGATCCTTACCCCTAAATATAAAAAGATTTCATAGAACATAAATCTCCAGAAAAAACATCGCCGAAGGGCATTA (SEQ ID NO:20)	869300
R117H	3	none	invader	AATCATAGCTTCCTATGACCCGGATAACAAGGAGGAACT (SEQ ID NO:21)	443800
R117H	3	hex	probe/DM	CGCGCCGAGGACTCTATCGCGATTTATCT (SEQ ID NO:22)	304200
R117H	3	none	synthetic target	ATGCCTAGATAAATCGCGATAGAGTGTTCCTCCTTGTTATCCGGGTCATAGGAAGCTATGATT (SEQ ID NO:23)	681700
R560T	1	none	invader	CATGAATGACATTTACAGCAAATGCTTGCTAGACCAATAATTAGTTATTCACT (SEQ ID NO:24)	595000
R560T	1	hex	probe/ER24	ACGGACGCGGAGGTTGCTAAAGAAATTCTTGCT (SEQ ID NO:25)	378100
R560T	1	none	synthetic target	CAACGAGCAAGAATTTCTTTAGCAACGTGAATAACTAATTATTGGTCTAGCAAGCATTTGCTGTAAATGTCATTCATGTAAAA (SEQ ID NO:26)	945400
3120+1G>A	2	none	invader	GCAATTTTGGATGACCTTCTGCCTCTTACCATATTTGACTTCATCCAGT (SEQ ID NO:27)	496000
3120+1G>A	2	hex	probe/DM	CGCGCCGAGGATATGTAAAAATAAGTACCGTTAA (SEQ ID NO:28)	397500
3120+1G>A	2	none	synthetic target	AGACATACTTAACGGTACTTATTTTTACATATCTGGATGAAGTCAAATATGGTAAGAGGCAGAAGGTCATCCAAAATTGCTATATC (SEQ ID NO:29)	984000
3659delC	2	none	invader	GAGAGTTGGCCATTCTTGTATGGTTTGGTTGACTTT (SEQ ID NO:30)	372300
3659delC	2	hex	probe/DM	CGCGCCGAGGGTAGGTTTACCTTCTGTTGG (SEQ ID NO:31)	302800
3659delC	2	none	synthetic target	CATGCCAACAGAAGGTAAACCTACAAGTCAACCAAACCATACAAGAATGGCCAACTCTC (SEQ ID NO:32)	679800
A455E	1	none	invader	CCTGAAAGATATTAATTTCAAGATAGAAAGAGGACAGTTGTTGGT (SEQ ID NO:33)	531000
A455E	1	hex	probe/ER24	ACGGACGCGGAGAGGTTGCTGGATCCA (SEQ ID NO:34)	298100
A455E	1	none	synthetic target	CCAGTGGATCCAGCAACCTCCAACAACTGTCCTCTTTCTATCTTGAAATTAATATCTTTCAGG (SEQ ID NO:35)	661000
1078delT	2	none	invader	AGTGCATAGGGAAGCACAGATAAAAACACCACAT (SEQ ID NO:36)	413500
1078delT	2	hex	probe/DM	CGCGCCGAGGAGAACCCTGAGAAGAAGAA (SEQ ID NO:37)	355400
1078delT	2	none	synthetic target	AGCCTTCTTCTTCTCAGGGTTCTTGTGGTGTTTTTATCTGTGCTTCCCTATGCACT (SEQ ID NO:38)	533300
G551D	2	none	invader	GCAGAGAAAGACAATATAGTTCTTGGAGAAGGTGGAATCACACTGAGTGGAGT (SEQ ID NO:39)	628200
G551D	2	hex	probe/DM	CGCGCCGAGGATCAACGAGCAAGAATTTCT (SEQ ID NO:40)	343800
G551D	2	none	synthetic target	CTTGCTAAAGAAATTCTTGCTCGTTGATCTCCACTCAGTGTGATTCCACCTTCTCCAAGAACTATATTGTCTTTCTCTGCAAACTT (SEQ ID NO:41)	883100
I148T	1	none	invader	AAATCAAACTAAACATAGCTATTCTCATCTGCATTCCAT (SEQ ID NO:42)	432400
I148T	1	hex	probe/ER24	ACGGACGCGGAGGTGTGATGAAGGCCAAA (SEQ ID NO:43)	350200
I148T	1	none	synthetic target	CCATTTTTGGCCTTCATCACACTGGAATGCAGATGAGAATAGCTATGTTTAGTTTGATTT (SEQ ID NO:44)	643100
N1303K	2	none	invader	CCATATTTCTTGATCACTCCACTGTTCATAGGGATCCAAT (SEQ ID NO:45)	414700
N1303K	2	hex	probe/DM	CGCGCCGAGGCTTTTTTCTAAATGTTCCAGAAAAA (SEQ ID NO:46)	391200
N1303K	2	none	synthetic target	ATTTATTTTTTCTGGAACATTTAGAAAAAAGTTGGATCCCTATGAACAGTGGAGTGATCAAGAAATATGGAAAG (SEQ ID NO:47)	867100
711+1G>T	2	none	invader	GCCTTTCCAGTTGTATAATTTATAACAATAGTGCCTAAAAGATTAAATCAATAGGTACATT (SEQ ID NO:48)	
711+1G>T	2	hex	probe/DM	CGCGCCGAGGAATTCATCAAATTTGTTCAGGT (SEQ ID NO:49)	
711+1G>T	2	none	synthetic target	ACCTGAACAAATTTGATGAATTATGTACCTATTGATTTAATCTTTTAGGCACTATTGTTATAAATTATACAACTGGAAAGGC (SEQ ID NO:50)	927000
1717-1G>A	3	none	invader	GCCTTTCAAATTCAGATTGAGCATACTAAAAGTGACTCTCTAATTTTCTATTTTTGGTAATAT (SEQ ID NO:51)	685000

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Figure 2 cont'd

1717-1G>A	3	hex	probe/DM	CGCGCCGAGGAGACATCTCCAAGTTTGC (SEQ ID NO:52)	294500
1717-1G>A	3	none	synthetic target	CTCTGCAAACTTGGAGATGTCTTATTACCAAAAATAGAAAATTAGAGAGTCACTTTTAGTATGCTCAATCTGAATTTGAAAGGCACATC (SEQ ID NO:53)	1010000
W1282X	3	none	invader	GCTCACCTGTGGTATCACTCCAAAGGCTTTCCTA (SEQ ID NO:54)	345000
W1282X	3	hex	probe/DM	CGCGCCGAGGTCACTGTTGCAAAGTTATTG (SEQ ID NO:55)	327800
W1282X	3	none	synthetic target	GATTCAATAACTTTGCAACAGTGAAGGAAAGCCTTTGGAGTGATACCACAGGTGAGCAA (SEQ ID NO:56)	683000
3849+10kbC>T	2	none	invader	CAAGAGTCTTCCATCTGTTGCAGTATTAAAATGGA (SEQ ID NO:57)	390000
3849+10kbC>T	2	hex	probe/DM	CGCGCCGAGGTGAGTAAGACACCCTGAAA (SEQ ID NO:58)	327400
3849+10kbC>T	2	none	synthetic target	TTCCTTTCAGGGTGTCTTACTCACCATTTTAATACTGCAACAGATGGAAGACTCTTG (SEQ ID NO:59)	601000
R553X	4	none	invader	CATTTACAGCAAATGCTTGCTAGACCAATAATTAGTTATTCACCTTGCTAAAGAAATTCTTGCTG (SEQ ID NO:60)	
R553X	4	hex	probe/DM	CGCGCCGAGGCATTGACCTCCACTCAGT (SEQ ID NO:61)	
R553X	4	none	synthetic target	ACTGAGTGGAGGTCAATGAGCAAGAATTTCTTTAGCAAGGTGAATAACTAATTATTGGTCTAGCAAGCATTTGCTGTAAATG (SEQ ID NO:62)	
G542X	4	none	invader	TCCAAGTTTGCAGAGAAAGACAATATAGTTCTTTC (SEQ ID NO:63)	
G542X	4	hex	probe/DM	CGCGCCGAGGGAGAAGGTGGAATCACA (SEQ ID NO:64)	
G542X	4	none	synthetic target	TGTGATTCCACCTTCTCAAAGAACTATATTGTCTTTCTCTGCAAACTTGGA (SEQ ID NO:65)	
621+1G>T	3	none	invader	CCTTCATCACATTGGAATGCAGATGAGAATAGCTATGTTTAGTTTGATTTATAAGAAGC (SEQ ID NO:66)	664000
621+1G>T	3	hex	probe/DM	CGCGCCGAGGTTAATACTTCCTTGCACAGG (SEQ ID NO:67)	311900
621+1G>T	3	none	synthetic target	GGGGCCTGTGCAAGGAAGTATTAACTTCTTATAAATCAAACTAAACATAGCTATTCTCATCTGCATTCCAATGTGATGAAGGCCAA (SEQ ID NO:68)	965000
R334W	2	none	invader	CGCAGAACAATGCAGAATGAGATGGTGGTGAATATTTTCCT (SEQ ID NO:69)	467000
R334W	2	hex	probe/DM	CGCGCCGAGGAGAGGATGATTCCTTTGATTA (SEQ ID NO:70)	336800
R334W	2	none	synthetic target	TGCACTAATCAAAGGAATCATCCTCTGGAAAATATTCACCACCATCTCATTCTGCATTGTTCTGCG (SEQ ID NO:71)	703000

Mutation	Exemplary Pool	3' blocking group	Oligo type (Arm)	Sequence (5'-3')	ε_{260} M^{-1} cm^{-1}
Internal control	all	none	Invader	tgtacttcatgctgtctacactaagagagaatgagagacacaca (SEQ ID NO:72)	503500
Internal control	all	hex	Probe/SNP4b	tccgcgcgtcctgaagaagcaccaatcatg (SEQ ID NO:73)	321200
Internal control	all	none	Synthetic Target	tttcatgattggtgcttcttcagtgtgtctctcattctctcttagtgtagacagcatgaagtacattt (SEQ ID NO:74)	698200

all	Hex	DM/FAM	Y-tct-X-agc-cgg-ttt-tcc-ggc-tga-gac-ctc-ggc-gcg-hex (SEQ ID NO:75)
1, 2, 5	Hex	ER24/FAM	Y-tct-X-agc-cgg-ttt-tcc-ggc-tga-gac-tcc-gcg-tcc-gt-hex (SEQ ID NO:76)
all	Hex	SNP4b/Red	Y-tct-X-tcg-gcc-ttt-tgg-ccg-aga-gag-gac-gcg-cgg-a-hex (SEQ ID NO:77)

X = Quencher = Z28
Y = Dye = FAM for 1055-48-08 and 1055-48-09 and Y = Z35 (or Redmond Red) for 1055-49-04

Mutation	Pool	3' blocking group	Oligo type (Arm)	Sequence (5'-3')	ε_{260} M^{-1} cm^{-1}
delF508	delF508	none	Invader	TGATGACGCTTCTGTATCTATATTCATCATAGGAAACACA (SEQ ID NO:78)	441500
delF508	delF508	Hex	WT Probe	CGCGCCGAGGCAAAGATGATATTTTCTTTAATGGT (SEQ ID NO:79)	382200
delF508	delF508	Hex	Mut Probe	AGCTCGTCCGACACAATAATATTTTCTTTAATGGTGCCA (SEQ ID NO:80)	418100
delF508	delF508	Hex	DM/FAM	Y-tct-X-agc-cgg-ttt-tcc-ggc-tga-gac-ctc-ggc-gcg-hex (SEQ ID NO:81)	347150
delF508	delF508	Hex	Wingra/Red	Y-tct-X-tcg-gcc-ttt-tgg-ccg-aga-gat-gtc-gga-cga-gct-hex (SEQ ID NO:82)	390400
delF508	delF508	none	WT Target	TGCCTGGCACCATTAAAGAAAATATCATCTTTGGTGTTTCCTATGATGAATATAGATACAGAAGCGTCATCAAA (SEQ ID NO:83)	837500
delF508	delF508	none	Mut Target	ATGCCTGGCACCATTAAAGAAAATATCATTGGTGTTTCCTATGATGAATATAGATACAGAAGCGTCATCAAA (SEQ ID NO:84)	828100

Mutation	Pool	3' blocking group	Oligo type (Arm)	Sequence (5'-3')
2184delA	2184delA	none	Invader	CTTCCTTTTTTCCCCAAACTCTCCAGTCTGTTTAAAAGATTGTTTA (SEQ ID NO:85)
2184delA	2184delA	hex	MUT probe/DM	CGCGCCGAGGTTTGTTTCTGTCCAGGAG (SEQ ID NO:86)
2184delA	2184delA	hex	WT probe/ER24	ACGGACGCGGAGTTTTGTTTCTGTCCAGGAG (SEQ ID NO:87)

X = Quencher = Z28
Y = Dye = FAM for 1055-48-08 and Y = Z35 (or Redmond Red) for 1144-16-02

711+1G>T	2	hex	probe/ER24	ACGGACGCGGAGAATTCATCAAATTTGTTCAGG (SEQ ID NO:88)
3849+10kb	2	hex	probe/ER24	ACGGACGCGGAGTGAGTAAGACACCCTGAAA (SEQ ID NO:89)

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